

SEQ LISTING\_ST25.txt  
SEQUENCE LISTING

<110> Frey, Joachim  
Stuber, Katja  
Thornton, Julian C  
Kuzyk, Michael A  
Burian, Jan

<120> TYPE III SECRETION PATHWAY IN AEROMONAS SALMONICIDA, AND USES THEREFOR

<130> VA/H-50095

<140> US 10/813,908  
<141> 2004-03-26

<150> US 10/416,902  
<151> 2003-05-15

<150> PCT/CA01/01589  
<151> 2001-11-15

<150> US 60/248,864  
<151> 2000-11-15

<160> 19

<170> PatentIn version 3.5

<210> 1  
<211> 47  
<212> PRT  
<213> Aeromonas salmonicida

<400> 1

Glu Leu Lys Arg Leu Ile Arg Leu Leu Pro Val Glu Leu Phe Ser Glu  
1 5 10 15

Glu Glu Gln Arg Gln Asn Leu Leu Gln Cys Cys Gln Gly Ala Leu Asp  
20 25 30

Asn Ala Ile Glu Arg Glu Glu Asp Glu Leu Ser Gly Glu Ser Ser  
35 40 45

<210> 2  
<211> 123  
<212> PRT  
<213> Aeromonas salmonicida

<400> 2

Met Asn Trp Ile Glu Pro Leu Leu Val Gln Phe Cys Gln Asp Leu Gly  
1 5 10 15

Ile Thr Ile Gly Asp Asn Pro His Ser Leu Ile Gln Leu Glu Leu Glu  
20 25 30

SEQ LISTING\_ST25.txt

Gln Ser Gly Thr Leu Gln Leu Glu Arg His Gln Gly Gln Leu Thr Leu  
35 40 45

Trp Leu Ala Arg Ala Val Pro Trp His Gln Ser Gly Glu Ala Ile Arg  
50 55 60

Arg Ala Met Thr Leu Thr Ala Ala Ala Gln Gly Pro Ala Leu Pro Val  
65 70 75 80

Arg Ser Gly Trp Leu Gly Glu Glu Gln Leu Ile Leu Phe Val Ser Leu  
85 90 95

Asp Glu Arg Ala Val Thr Leu Pro Gln Leu His Gln Ala Val Thr Thr  
100 105 110

Leu Thr Arg Leu Gln Arg Glu Val Leu Ala Ser  
115 120

<210> 3  
<211> 121  
<212> PRT  
<213> Aeromonas salmonicida

<400> 3

Met Ser Arg Ile Thr Ala Ala His Ile Gly Ile Glu Gln Leu Ser Ala  
1 5 10 15

Ile Ser Leu Asp Asp Gln Glu Arg Ser Leu Pro Gly Arg Tyr Ala Leu  
20 25 30

Leu Pro Asp Gly Gln Ser Ile Glu Pro His Ile Ser Arg Leu Tyr Pro  
35 40 45

Glu Arg Leu Ala Asp Arg Val Leu Leu Asp Phe Ala Thr Pro Asp Arg  
50 55 60

Gly Phe His Asp Leu Leu Arg Pro Val Asp Phe Asn Gln Ala Met Gln  
65 70 75 80

Gly Leu Arg Ser Val Leu Ala Glu Gly Gln Ser Pro Glu Leu Arg Ala  
85 90 95

Ala Ala Ala Leu Leu Glu Gln Met His Ala Asp Glu Gln Leu Met Gln  
100 105 110

Met Thr Leu His Leu Leu His Lys Val  
115 120

SEQ LISTING\_ST25.txt

<210> 4  
 <211> 116  
 <212> PRT  
 <213> Aeromonas salmonicida  
 <400> 4  
 Met Thr Met Val Leu Thr Ser Gln Gln Gln Asp Ala Leu Leu Leu Thr  
 1 5 10 15  
 Gly Trp Leu Gln Leu Gln Tyr Gly His Pro Asp Lys Ala Ser Val Leu  
 20 25 30  
 Leu Ala Ala Leu Leu Gln Ile His Pro Asp His Gln Gly Gly Arg Arg  
 35 40 45  
 Thr Leu Leu Val Ala Leu Leu Lys Gln Gly Glu Gly Glu Ala Ala Leu  
 50 55 60  
 Ala His Val Asp Gln Leu Met Gln Gln Gly Glu Ala Asp Gly Pro Leu  
 65 70 75 80  
 Trp Leu Cys Arg Ser Arg Ala Cys Gln Leu Ala Gly Arg Leu Asp Glu  
 85 90 95  
 Ala Arg Phe Ala Tyr Gln Gln Tyr Leu Glu Leu Glu Glu Gln Asn Glu  
 100 105 110  
 Ser Thr His Pro  
 115

<210> 5  
 <211> 705  
 <212> PRT  
 <213> Aeromonas salmonicida  
 <400> 5  
 Met Asn Gln Arg Thr Leu Glu Leu Leu Arg Arg Ile Gly Glu Arg Lys  
 1 5 10 15  
 Asp Ile Met Leu Ala Ile Leu Leu Leu Ala Ile Val Phe Met Met Val  
 20 25 30  
 Leu Pro Leu Pro Pro Val Ala Leu Asp Ile Leu Ile Ala Ile Asn Met  
 35 40 45  
 Thr Ile Ser Val Val Leu Leu Met Met Ala Val Tyr Ile Asn Ser Pro  
 50 55 60  
 Leu Gln Phe Ser Ala Phe Pro Ala Val Leu Leu Ile Thr Thr Leu Phe  
 Page 3

SEQ LISTING\_ST25.txt

65		70		75		80									
Arg	Leu	Ala	Leu	Ser	Val	Ser	Thr	Thr	Arg	Met	Ile	Leu	Leu	Gln	Ala
				85					90					95	
Asp	Ala	Gly	Gln	Ile	Val	Tyr	Thr	Phe	Gly	Asn	Phe	Val	Val	Gly	Gly
			100					105					110		
Asn	Leu	Val	Val	Gly	Ile	Val	Ile	Phe	Leu	Ile	Ile	Thr	Ile	Val	Gln
		115					120					125			
Phe	Leu	Val	Ile	Thr	Lys	Gly	Ser	Glu	Arg	Val	Ala	Glu	Val	Ser	Ala
	130					135					140				
Arg	Phe	Ser	Leu	Asp	Ala	Met	Pro	Gly	Lys	Gln	Met	Ser	Ile	Asp	Gly
145					150					155				160	
Asp	Met	Arg	Ala	Gly	Val	Ile	Asp	Val	His	Glu	Ala	Arg	Asp	Arg	Arg
				165					170				175		
Gly	Val	Ile	Glu	Lys	Glu	Ser	Gln	Met	Phe	Gly	Ser	Met	Asp	Gly	Ala
		180						185					190		
Met	Lys	Phe	Val	Lys	Gly	Asp	Ala	Ile	Ala	Gly	Leu	Ile	Ile	Ile	Phe
		195					200					205			
Val	Asn	Ile	Leu	Gly	Gly	Val	Thr	Ile	Gly	Val	Thr	Gln	Lys	Gly	Leu
	210					215					220				
Ser	Ala	Ala	Asp	Ala	Leu	Gln	Leu	Tyr	Ser	Ile	Leu	Thr	Val	Gly	Asp
225					230					235				240	
Gly	Met	Val	Ser	Gln	Val	Pro	Ala	Leu	Leu	Ile	Ala	Ile	Thr	Ala	Gly
				245					250					255	
Ile	Ile	Val	Thr	Arg	Val	Ser	Ser	Glu	Glu	Ser	Ser	Asp	Leu	Gly	Thr
		260						265					270		
Asp	Ile	Gly	Ala	Gln	Val	Val	Ala	Gln	Pro	Lys	Ala	Leu	Leu	Ile	Gly
	275						280					285			
Gly	Leu	Leu	Leu	Val	Leu	Phe	Gly	Leu	Ile	Pro	Gly	Phe	Pro	Met	Ile
	290					295					300				
Thr	Phe	Phe	Ala	Leu	Ser	Ala	Ile	Val	Thr	Ala	Gly	Gly	Tyr	Phe	Ile
305					310					315				320	

SEQ LISTING\_ST25.txt

Gly Leu Arg Gln Arg Lys Ala Gln Ser Ser Asn Ser Gln Asp Leu Pro  
325 330 335

Ala Val Leu Ala Gln Gly Ala Gly Ala Pro Ala Ala Arg Ser Lys Pro  
340 345 350

Lys Pro Gly Ser Lys Pro Arg Gly Lys Leu Gly Glu Lys Glu Glu Phe  
355 360 365

Ala Met Thr Val Pro Leu Leu Ile Asp Val Asp Ala Ala Leu Gln Ala  
370 375 380

Glu Leu Glu Ala Ile Ala Leu Asn Asp Glu Leu Val Arg Val Arg Arg  
385 390 395 400

Ala Leu Tyr Leu Asp Leu Gly Val Pro Phe Pro Gly Ile His Leu Arg  
405 410 415

Phe Asn Glu Gly Met Gly Pro Gly Glu Tyr Leu Ile Gln Leu Gln Glu  
420 425 430

Val Pro Val Ala Arg Gly Leu Leu Arg Pro Gly His Gln Leu Val Gln  
435 440 445

Glu Ser Ala Ser Gln Leu Asp Leu Leu Gly Ile Pro Tyr Glu Glu Gly  
450 455 460

Ala Pro Leu Leu Pro Gly Gln Pro Thr Leu Trp Val Ala Asn Glu His  
465 470 475 480

Gln Glu Arg Leu Glu Lys Ser Arg Leu Ala Thr Leu Thr Thr Asp Gln  
485 490 495

Val Met Thr Trp His Leu Ser His Val Leu Arg Glu Tyr Ala Glu Asp  
500 505 510

Phe Ile Gly Ile Gln Glu Thr Arg Tyr Leu Leu Glu Gln Met Glu Gly  
515 520 525

Ser Tyr Ser Glu Leu Val Lys Glu Ala Gln Arg Ile Ile Pro Leu Gln  
530 535 540

Arg Met Thr Glu Ile Leu Gln Arg Leu Val Gly Glu Asp Ile Ser Ile  
545 550 555 560

Arg Asn Met Arg Ala Ile Leu Glu Ala Met Val Glu Trp Gly Gln Lys  
565 570 575

SEQ LISTING\_ST25.txt

Glu Lys Asp Val Val Gln Leu Thr Glu Tyr Ile Arg Ser Ser Leu Lys  
580 585 590

Arg Tyr Ile Cys Tyr Lys Tyr Ala Asn Gly Asn Asn Ile Leu Pro Ala  
595 600 605

Tyr Leu Leu Asp Gln Gln Val Glu Glu Gln Leu Arg Gly Gly Ile Arg  
610 615 620

Gln Thr Ser Ala Gly Ser Tyr Leu Ala Leu Asp Pro Thr Ile Thr Gln  
625 630 635 640

Ser Phe Leu Asp Gln Val Arg His Thr Val Gly Asp Leu Ala Gln Met  
645 650 655

Gln Asn Lys Pro Val Leu Ile Val Ser Met Asp Ile Arg Arg Tyr Val  
660 665 670

Arg Lys Leu Ile Glu Gly Asp Tyr His Ala Leu Pro Val Leu Ser Tyr  
675 680 685

Gln Glu Leu Thr Gln Gln Ile Asn Ile Gln Pro Leu Gly Arg Val Cys  
690 695 700

Leu  
705

<210> 6  
<211> 93  
<212> PRT  
<213> Aeromonas salmonicida  
  
<400> 6

Met Leu Val Arg Arg Glu Gly Glu Arg Ala Gly Leu Ala Asn Pro Phe  
1 5 10 15

Ala Ala Leu Tyr Leu Leu Ala Glu Ala Thr Leu Ala Val Leu Gly Pro  
20 25 30

Gly His Phe Leu Tyr Gly Asn Val Asp Val Phe Arg Ser Ser Ser Leu  
35 40 45

Ser Ser Glu Arg Leu Gly Arg Phe Tyr Leu Arg Trp Thr Gly Ala Ser  
50 55 60

Glu Pro Glu Pro Gly Trp Phe Met Leu Ala Thr Glu Gln Val Cys Ser  
65 70 75 80

SEQ LISTING\_ST25.txt

Leu Arg Asp Met Arg Lys Arg Gln Lys His Gly Leu Ala  
85 90

<210> 7  
<211> 94  
<212> PRT  
<213> Aeromonas salmonicida

<400> 7

Met Lys Gln Pro Arg Phe Ala Asp His Ser Glu Thr Ile Ser Gln Ala  
1 5 10 15

Glu His Gly Ile Ala Asp Ser Asp His Arg Asn Ala Leu Leu Gln Glu  
20 25 30

Met Leu Ala Gly Leu Ala Leu Ser Asp Gln Thr Cys Gln Leu Leu Phe  
35 40 45

Glu Ala Pro Thr Glu Gln Val Ala Val Ala Glu Gln Glu Leu Leu Ala  
50 55 60

Glu Ile Gln Arg Arg Gln Ala Leu Leu Pro Ala Gln Pro Gly Glu Gly  
65 70 75 80

Arg Lys Ser Arg Arg Pro Thr Ile Met Arg Gly Leu Met Ile  
85 90

<210> 8  
<211> 361  
<212> PRT  
<213> Aeromonas salmonicida

<400> 8

Met Ser Thr Ile Pro Asp Tyr Asn Thr Asn Pro Gly Ala Phe Val Gly  
1 5 10 15

Trp Leu Asp Val Gln Ala Leu Asn Thr Leu Pro Gly Asn Lys Asn Pro  
20 25 30

Lys Leu Thr Glu Leu Val Glu Leu Leu Lys Gly Lys Ile Thr Ile Ser  
35 40 45

Ala Asp Ser Ser Thr Ala Leu Ser Lys Glu Gln Leu Glu Lys Leu Leu  
50 55 60

Ala Ala Tyr Leu Thr Asp Pro Ala Ser Ile Asn Gly Gly Trp Ala Met  
65 70 75 80

SEQ LISTING\_ST25.txt

Gly Gln Phe Lys Gly Gly Gln Asp Ala Ala Ile Ala Ala Ile Lys Gly  
85 90

Val Ile Glu Arg Gly Ala Lys Gln Thr Pro Pro Val Thr His Trp Thr  
100 105 110

Ile Pro Glu Phe Met Leu Leu Ser Leu Ser Ala Leu Thr Met Glu Arg  
115 120 125

Thr Asp Asp Asp Leu Ile Thr Thr Phe Thr Gly Val Met Met Phe Gln  
130 135 140

Asp Asn Gln Arg Lys Gly Leu Arg Asp Glu Leu Ala Glu Met Thr Ala  
145 150 155 160

Glu Leu Lys Ile Tyr Gly Val Ile Gln Ser Glu Ile Asn Gln Val Leu  
165 170 175

Ser Ala Ala Ser Asn Gln Thr Phe Lys Thr Asn Phe Asn Leu Met Asp  
180 185 190

Tyr Lys Leu Tyr Gly Tyr Glu Ser Leu Ala Lys Phe Met Glu Gly Gly  
195 200 205

Glu Phe Lys Leu Leu Ser Lys Met Phe Ser Asp Glu Gln Val Thr Lys  
210 215 220

Ala Gln Gln Asp Phe Thr Asn Ala Lys Asn Glu Leu Glu Asn Val Thr  
225 230 235 240

Ser Thr Ser Leu Asn Pro Lys Ile Gln Ala Glu Ala Lys Thr Asp Tyr  
245 250 255

Glu Arg Lys Lys Ala Ile Phe Glu Glu Ile Val Glu Thr Gln Ile Ile  
260 265 270

Thr Leu Lys Thr Phe Leu Glu Ser Asp Leu Lys Lys Ser Gly Ala Met  
275 280 285

Ser Gly Ile Glu Ala Glu Tyr Lys Tyr Asp Lys Asp Asn Asn Lys Leu  
290 295 300

Gly Asn Phe Ser Thr Ser Val Ser Asp Arg Ser Arg Pro Leu Asn Asp  
305 310 315 320

Leu Val Ser Glu Lys Thr Ala Arg Leu Asn Asp Val Ser Ser Arg Tyr  
325 330 335



SEQ LISTING\_ST25.txt

Asn Ala Ala Ile Glu Ala Leu Asn Arg Phe Ile Gln Lys Tyr Asp Ser  
340 345 350

Ile Met Arg Asp Ile Leu Gly Ala Ile  
355 360

<210> 9  
<211> 159  
<212> PRT  
<213> Aeromonas salmonicida

<400> 9

Met Gln Thr Asp Thr Thr Leu Thr Pro Glu Tyr Glu Ala Glu Leu Glu  
1 5 10 15

Ala Phe Met Ala Asp Gly Gly Thr Leu Ala Met Leu Gln Asp Ile Ser  
20 25 30

Gly Asp Thr Leu Glu Gln Leu Tyr Ala Leu Ala Phe Ser Gln Tyr Gln  
35 40 45

Ala Gly Lys Trp Glu Asp Ala His Lys Ile Phe Gln Ala Leu Cys Met  
50 55 60

Leu Asp His Tyr Glu Pro Arg Tyr Phe Leu Gly Leu Gly Ala Cys Arg  
65 70 75 80

Gln Ala Met Gly Glu Phe Glu Thr Ala Val Gln Ser Tyr Ser Phe Gly  
85 90 95

Ala Met Leu Asp Leu Lys Asp Pro Arg Phe Pro Phe His Ala Gly Glu  
100 105 110

Cys Arg Leu Gln Gln Gly Asp Leu Asn Gly Ala Glu Ser Gly Phe His  
115 120 125

Ser Ala Arg Leu Leu Ala Asp Thr Asp Pro Gln Gln Ala Asp Leu Ala  
130 135 140

Ala Ser Ala Lys Val Met Leu Glu Ala Ile Ala Ile Arg Arg Asp  
145 150 155

<210> 10  
<211> 5678  
<212> DNA  
<213> Aeromonas salmonicida

<400> 10

gagctcaagc ggctgatccg cctgctgccg gtggagctgt tcagtgaga ggagcagcg

60

SEQ LISTING\_ST25.txt

cagaatctgt tgcagtgtg tgcaggtgctg ctcgataacg ccatcgagcg ggaagaggat	120
gagttgtctg gagagtcgtc atgaactgga ttgaacccct gctggtgcag ttttgccagg	180
atttgggcat caccataggg gataaccccc attcgtgat ccagcttgaa ctggagcaga	240
gcggcactct gcagctggag cgccatcagg ggcaactgac cctatggttg gcccgcgccg	300
tgccctggca tcagagtggc gaggccattc gcccgccat gacctgact gcccgggcgc	360
aagggccggc actgccggtg cgcagcggtt ggttggggga ggagcagttg atctctttcg	420
tctcctgga tgagcgggccc gtgactctgc cccagctcca tcaggccgtg accacctga	480
cccggttgca gcgagaggtg ctggcgctcat gagccggatc actgccgcgc atatcggtat	540
cgagcagctc agcgccatct ccctcgacga tcaggagcgc agcctgccgc ggcgttatgc	600
cctgttgccc gatggccagt ccatcgaacc ccatatcagc cgcctctacc ccgagcggt	660
ggcggatcgg gtgctgctcg atttcgccac cccggatcgc ggctttcacg acttgctgcg	720
accggtcgat ttcaatcagg cgatgcaggg gctgcgcagt gtgctggcag aggggcagag	780
ccccgaattg cgagcgggccc ccgcgctgct cgaacaaatg cacgccgatg aacaactgat	840
gcagatgacc cttcatctgc tgcacaaggt atgacctgg tgcttacgtc acagcagcag	900
gatgcgctgc tgctcaccgg ctggttgcaa ctgcaatatg gccacctga caaggcgagc	960
gtgctgctgg ccgccctgct gcagatccac cccgaccatc agggaggggc acggaccttg	1020
ctggtggccc tgctcaaaca gggggagggg gaggcggcgc tggcccatgt cgatcagctg	1080
atgcagcaag gggaggccga cggcccgctc tggctctgtc gcagccgagc ctgcagttg	1140
gcaggcgggc tggatgaagc ccgttttgcc tatcaacaat acctcgaact ggaagagcag	1200
aatgaatcaa cgcacccttg agttgctgcg ccggataggc gaacgaagg acatcatgct	1260
ggcgatcctg ctgctggcca tcgtctttat gatggtcttg ccgctgccgc cggtgccct	1320
cgatatcctg attgccatca acatgaccat ctcggttgta ctgctgatga tggcggttta	1380
tatcaattcg ccgctgcagt tctccgcctt tccggcggtg ctgctgatca ccacctgtt	1440
ccgcttgccc ttgctgggta gtaccacccg gatgatcctg ctgcaggctg atgcggggca	1500
gatagtctac accttcggca acttcgtggt ggggggcaat ctggtggtgg ggatcgctat	1560
cttctctatc atcaccatcg tccagtttct ggtgatcacc aagggctcgc agcgggtcgc	1620
cgaggtgagc gccgcgtttt ccctcgatgc catgccgggt aagcagatga gtatcgatgg	1680
tgacatgccc gccgggggta tcgacgtgca cgaggcgcg gatcgccgcg gggtcatcga	1740
gaaggagagc catagtttcg gctccatgga tggcgccatg aagtttgta agggggacgc	1800
catcgcgggc ctcacatca tcttcgtcaa ctcctcgtt ggctcacca tcgggggtgac	1860
ccagaagggt ttatccgccg ccgatgcgct gcagctctac tccatcctga cggtggtgta	1920

SEQ LISTING\_ST25.txt

tggcatggtc	tcccaggtgc	cggcgctgct	gatcgccatc	accgcgggca	tatatctcac	1980
ccgggtctcc	tccgaagagt	cttccgatct	gggtaccgat	atcggcgccc	aggtggtggc	2040
ccagccaag	gcgctactga	tcggcggtct	gctgctggtg	ctgttcgggt	tgatccccgg	2100
cttcccgatg	atcaccttct	ttgcgctgtc	ggccatcgct	acggcgggcg	gttactttat	2160
cggcttgca	caacgcgaag	cgcaaacgag	caacagtcag	gatcttctcg	ccgtgctggc	2220
gcagggggcc	ggggcccccag	ctgcccgcag	caagccaaaa	ccgggcagca	agccgcgggg	2280
caagctgggg	gagaaggagg	agtttgccat	gacggtgccg	ctccttatcg	atgtggatgc	2340
tgctttgcag	gccgagctgg	aggcgattgc	cctcaacgac	gaactggtgc	gggtgcgccg	2400
cgccctctat	ctcgatctcg	gggtgccttt	cccgggtatt	cacctgcgtt	tcaacgaggg	2460
gatggggcct	ggcgaatacc	tgatccagct	gcaggagggtg	ccggtcgccc	gcggtctgct	2520
gcgcccgggc	catcagctgg	tgacaggagag	cgctctccag	ctcgatctgc	tggggatccc	2580
ctacgaagag	ggggcgccgt	tactgccggg	acaaccgacc	ttgtgggtcg	ctaatagaaca	2640
tcaggagcga	ctggagaagt	cacggctggc	caccctcacc	accgatcagg	tgatgacctg	2700
gcacttatcc	catgtgctgc	gggaatatgc	cgaggacttt	atcggcattc	aggagacccg	2760
ctacctgctg	gagcagatgg	aggggagcta	tagcgagctg	gtgaaggagg	cgcaacgcac	2820
catcccgcgt	cagcgtatga	cgaaaatttt	gcagcggctg	gtgggggagg	atatctccat	2880
ccgcaacatg	cgcgccatcc	tcgaggcgat	ggtggagtgg	ggccagaagg	agaaggatgt	2940
ggtgcagctc	accgagtaca	tccgtagcag	cctcaagcgc	tacatctgct	acaagtacgc	3000
caacggcaac	aacattttgc	ctgcctatct	gctcgatcag	caggtggagg	agcagctccg	3060
cggcggcatt	cgccagacta	gtgccggcag	ctatctggcg	ctcgatccca	ctattaccca	3120
gagcttcctc	gatacaggtg	gccacaccgt	cggatgatctg	gcccagatgc	agaacaaacc	3180
ggtgtctcatt	gtctccatgg	atatccgccg	ctatgtgcgc	aagctcatcg	aggggggatta	3240
ccatgccctg	ccggtgctct	cctatcaggga	gctgaccacg	cagatcaata	tccagccctc	3300
cgggagggtg	tgccctgtgag	gggggaccgg	ttaacctctg	acccccgat	cccttggtcg	3360
caggccaagg	gtgtggcggt	tgccctctcac	tatctggggg	caacccccat	ccagctgggc	3420
cacgctttct	gctatcgcca	aatttatctc	gcctggcggg	ttgatcctac	gaccgcagcg	3480
gtctggtatg	tgctggtgcg	ccgagagggg	gagcgggctg	gactggccaa	tccctttgcc	3540
gccctctatc	tgctggccga	agccactctg	gctgtactcg	gtccgggcca	ttctctctac	3600
ggcaacgtcg	atgtctttcg	aagcagtagc	ctgagcagtg	agcggctagg	cgccttctac	3660
ttgcgctgga	cgggagccag	tgaacccgag	cccggctggt	tcattgtggc	caccgagcaa	3720
gtctgttcac	tacgggatat	gcgaaaacga	caaaagcacg	gccttgctgct	acaggcatgt	3780
ccaaaagggc	ctcatagaat	aggagccaag	atgaacaac	cgcgttttgc	cgaccatagc	3840

SEQ LISTING\_ST25.txt

gagaccattt cgcaggcaga gcatggcatt gccgacagcg atcaccgcaa tgccctgttg	3900
caagagatgc tggctggcct agccctctcg gatcagacct gtcagctgct gttcgaagcg	3960
ccgaccgagc aagtggccgt ggcgagcag gagtgttg cagagatcca gcgcagacag	4020
gcgttactac cggcacagcc gggagagggc cgcaaaagtc gccgtccac cattatgcgc	4080
ggactgatga ttttaaggagt cgtgatgagc acaatccctg actacaacac taacccggc	4140
gcgttcgtcg gctggcttga tgtgaagca ctgaacacat tgccgggcaa taaaatccc	4200
aagttgaccg aactggtcga gctgctcaag ggcaagatca ccatcagtgc tgactcatcg	4260
actgcgtga gcaaggagca gctggagaag ttgctggctg cctatctgac ggatcctgcc	4320
tcgatcaacg ggggctggg gatgggcca ttcaagggag gtcaagatgc gcgcattgcc	4380
gccatcaagg ggggtgatga gcggggagca aaacaaacc cgccagtcac ccactggacc	4440
atccctgaat ttatgctgct ctccctcagt gcgctgacca tggaaagtc cgatgacgat	4500
ctcatcacga cttttaccgg ggtgatgatg ttccaggaca atcagcgtaa aggggtgcg	4560
gatgagctgg cagagatgac cgctgagctg aagatctacg ggggatcca gtccgagatc	4620
aaccaggtgc tctctcggc gtccaacaa accttcaaaa ccaatttcaa tctgatggat	4680
tacaagctct atggctatga gtctctggcc aaatttatgg aaggggcgga gttcaagctg	4740
ttgtcaaaaa tgtttagcga tgagcagggt acaaaagcac agcaagattt caccaatgct	4800
aaaaatgagc tggaaaacgt cacgtcgacc agcctaaacc ccaaaatcca ggcggaagct	4860
aagaccgatt atgagcgtaa aaaagccatt tttgaggaga tcgtagagac gcagatcatc	4920
acccttaaaa cgttcctgga aagtgcctg aagaagagcg gcgccatgag tggcatagaa	4980
gccgagtaca aatatgacaa agacaacaac aagcttgga acttctccac tagtgtgagc	5040
gaccgttctc gcccgctcaa cgatctggtc agtgaaaaga cggccgcct caacgacgtc	5100
agttcgcgct acaacgctgc catcgaggca ctcaaccgct ttatccagaa atacgacagc	5160
atcatcgcg acattcttgg cgcaatttga ggagagatca tgcagaccga caccaccctg	5220
accccggaat atgaagcaga gctggaggcc tttatggctg acggtgtgac cctgctatg	5280
ctgcaggata tctctggcga caccttggaa cagctctatg ccctggcctt tagccagtat	5340
caggccggga agtggaaga tgctcacaaa atcttccagg ctctctgcat gctggatcac	5400
tacgagccac gctatttctt cgggctgggt gcttgccgtc aggcgatggg ggagtgtgaa	5460
acggcagttc agagttacag ctttggcgcc atgctcgacc tgaaagatcc ccgtttccca	5520
tttcatgcag cgagtgccg gctgcaacaa ggtgatttga acggtgccga gagtggcttc	5580
cactcggccc gactgctggc ggacacagat cccagcagg cagacctggc ggcaagcgcc	5640
aaggctcatg ttggaagcat cgcaatcaga agggatcc	5678

SEQ LISTING\_ST25.txt

<210>	11	
<211>	30	
<212>	DNA	
<213>	Artificial sequence	
<220>		
<223>	synthetic primer	
<400>	11	
	gggaattcga tgagcacaat ccctgactac	30
<210>	12	
<211>	30	
<212>	DNA	
<213>	Artificial sequence	
<220>		
<223>	Synthetic sequence	
<400>	12	
	atgcggccgc aaattgcgcc aagaatgtcg	30
<210>	13	
<211>	29	
<212>	DNA	
<213>	Artificial sequence	
<220>		
<223>	Synthetic primer	
<400>	13	
	tcgcggccgc accctttacg ctgattgtc	29
<210>	14	
<211>	28	
<212>	DNA	
<213>	Artificial sequence	
<220>		
<223>	Synthetic primer	
<400>	14	
	cgggaattcgt tgcgggatga gctggcag	28
<210>	15	
<211>	30	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	synthetic primer	
<400>	15	
	tcgcggccgc actcggcttc tatgccactc	30
<210>	16	
<211>	18	

SEQ LISTING\_ST25.txt

<212> DNA  
<213> Artificial sequence

<220>  
<223> Synthetic primer

<400> 16  
gcccgttttg cctatcaa 18

<210> 17  
<211> 18  
<212> DNA  
<213> Artificial sequence

<220>  
<223> Synthetic primer

<400> 17  
gcgccgatat cggtaccc 18

<210> 18  
<211> 20  
<212> DNA  
<213> Artificial sequence

<220>  
<223> Synthetic primer

<400> 18  
ttcgtcggct ggcttgatgt 20

<210> 19  
<211> 20  
<212> DNA  
<213> Artificial sequence

<220>  
<223> Synthetic primer

<400> 19  
gaactcgccc ccttcataa 20